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Global disparities in cystic fibrosis outcomes prior to CFTR modulators: A CF registries cohort study in South Africa and Canada

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ABSTRACT

Background: Outcomes of cystic fibrosis (CF) differ between low-middle income and high-income countries, but comparative data are lacking. We compared South African (SA) and Canadian CF outcomes to explore what disparities existed prior to access of CFTR modulators in Canada.

Methods: A cross-sectional study of SA and Canadian CF registries data for period 1 January to 31 December 2018. CF registry data were harmonised between countries to compare lung function and nutrition outcomes. Poor nutrition was defined as BMIz-score < -1 in children and < 18.5 kg/m² in adults. Standardised mean difference (SMD) >10 was considered significant.

Results: After excluding Canadians on CFTR modulators and lung transplant recipients, data on 4049 Canadian and 446 SA people was analysed. Compared to Canada, people in SA were younger (median age 15.8 years vs. 24.1 years; SMD 52) with fewer males (47.8% vs 54.2%; SMD 12.5) and White (70.9% vs. 93.3%; SMD 61.3). Class I-III CFTR mutation frequency was similar in SA ($n = 384$, 86.1%) and Canada ($n = 3426$, 84.9%). After adjusting for age, gender, diagnosis age, genotype, *P.aeruginosa* infection and pulmonary treatments, FEV1pp was 8.9% lower (95% CI 6.3% to 11.4%) and poor nutrition 1.7-fold more common (OR 1.70; 95% CI 1.19–2.41) in SA compared to Canada.

Conclusion: Lung function and nutrition was significantly lower in SA compared to Canada. Global disparities in CF outcomes between high and low-middle income countries are likely to widen as CFTR modulators are rapidly scaled up in only high-income countries.

1. Introduction

International cystic fibrosis (CF) registries from high income countries (HIC) have documented improving CF survival estimates over past decades [1]. Improving survival age is attributed to many factors including newborn screening (NBS), improved nutritional interventions, active surveillance, prevention and treatment of infections and organ transplantation [2]. In addition, novel cystic fibrosis transmembrane conductance regulator (CFTR) modulator drug therapy is anticipated to significantly further improve CF outcomes and survival [3,4]. In contrast

to HIC, less is known about the epidemiology and outcomes of CF populations in low-and middle-income countries (LMIC) where conditions of poverty, poor healthcare infrastructure and limited resources to diagnose CF prevail. Furthermore, ethnicity and CF genotype differ significantly in LMIC compared to HIC with populations of predominantly European descent [5,6]. Studies reporting CFTR mutation prevalence in Brazil and Africa have described higher rates of uncommon or novel mutations than HIC which may be an important determinant of CF outcomes in these settings [6,7]. However, factors such as socioeconomic status (SES), living conditions, delayed CF diagnosis, nutrition,

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pulmonary infections, and limited access to expensive CF therapies may be more important determinates of CF outcomes in LMIC compared to HIC.

South Africa (SA) is a middle income country with a population of nearly 60 million with diverse ancestry and conditions of extreme socioeconomic disparity [8]. Healthcare infrastructure and services are provided through a resource-constrained public health system and well-resourced private healthcare. Newborn screening for CF is not available. Essential CF care and CF medications including pancreatic enzyme replacement therapy (PERT) are available, but expensive CF therapies such inhaled tobramycin solution, recombinant DNase are scarce and CFTR modulator therapies are not available. Poor nutrition, methicillin resistant staphylococcus aureus (MRSA) infection and non-White ancestry have all been associated with poorer CF outcomes in SA [9,10]. In contrast, Canada is a HIC with a population of almost 40 million. Universal healthcare including NBS for CF is provided for free which allows Canadians to access most CF medical care without additional personal financial costs. The majority of essential CF medications and lung transplants are paid for by government programs including the early CFTR modulator drugs (ivacaftor), prior to 2018 for a small number of people with CF (pwCF) who were eligible.

Examining differences in health outcomes between CF care centers has spawned a body of evidence focused on quality improvement initiatives in an effort to explain such differences and improve outcomes [11,1]. Similarly, international comparisons of the CF population through registries have led to important research to understand the driving factors contributing to disparate CF outcomes [12,13]. Comparing current CF health outcomes in SA and Canada, and factors that potentially explain any difference in outcomes, may be helpful for planning targeted interventions and advocating for improved access to novel therapies where outcomes are less favourable. We therefore harmonised data in the SA and Canadian CF registries to compare demographic, clinical and health outcomes characteristics between people with pwCF receiving care in SA and Canada at a time prior to when CFTR modulator therapy was more widely available in Canada.

2. Methods

2.1. Study design and populations

This is a cross-sectional cohort study of CF populations captured in the SA and Canadian CF registries for the calendar period 1 January to 31 December 2018. The Canadian CF Registry (CCFR) was developed in the 1970s and is managed by Cystic Fibrosis Canada. Consented data are entered into the CCFR by the 42 CF centers across Canada. Universal access to health care and incentivised data entry programs for participating CF centers within Canada ensures that the majority of pwCF are captured in the CCFR. The SA CF Registry (SACFR) is a public-private collaboration that was launched in 2018 and includes CF care centers at six university hospitals and several private practice clinics across SA with expertise in CF care. Consented annual review registry data is collected by independent data capturers covering different regions of SA, who extract and capture registry data from medical records at each participating clinic or practice. By 31 December 2018, it is estimated that over 80% of the known SA CF population in care was captured in the SA CF registry [9].

2.2. Harmonization of CF registry data

A combined data dictionary outlining variable definitions was systematically created using a mapping strategy in order to make data directly comparable (see on-line supplement). Demographic, CF diagnosis and genotype information were collected. Genotype was classified as homozygous F508del, heterozygous F508del, not F508del and missing. CFTR mutations were further categorised as class I–III and class IV–VI. Clinical information in 2018 on sputum microbiology, CF-related

complications, CF therapies including number of courses of intravenous antibiotics, were recorded from both countries. Due to inability to harmonize most sputum microbiology data captured in the registries as a result of differing data collection definitions, sputum microbiology reporting was restricted to any isolate of *P.aeruginosa*, methicillin resistant *staphylococcus aureus* (MRSA) and non-tuberculous *mycobacterium species* (NTM).

2.3. Outcomes

Lung function (LF) was measured by forced expiratory volume in one second (FEV₁) expressed as a percentage of the predicted values (FEV₁pp) for healthy age, race and sex-matched controls using Global Lung Initiative (GLI) reference equations [14]. Non-White people in SA were entered as “Other” ethnicity in GLI equations in line with published guidelines [15]. In order to harmonize LF measurements, the highest recorded stable pre-bronchodilator FEV₁pp in 2018 for subjects six years and older were selected from the SA and Canadian registries as this is how FEV₁pp is captured in the SA CF registry. Lung function impairment was classified as severe (FEV₁pp ≤ 40), moderate (FEV₁pp 41–69), mild (FEV₁pp 70–89) and normal (FEV₁pp ≥ 90). In the event of death, lung transplantation or starting CFTR modulator therapy in 2018, patients were included but the FEV₁pp measurements in 2018 prior to these events were included in analyses. People who started CFTR modulators, died or were transplanted prior to 2018 were excluded from formal analyses (Fig. 1).

Body mass index (BMI) measurements were selected from the same day as FEV₁ measurements. BMI z-scores (BMIZ) were calculated for children 0–18 years using World Health Organization reference equations [16]. For individuals 18 years of age and older, BMI (kg/m²) was calculated and classified according to World Health Organization guidelines. Poor nutrition was defined as BMIZ < -1 for age < 18 years or BMI < 18.5 kg/m² for age 18 years and older; overweight was defined as BMIZ > 2 for age < 18 years or BMI > 24.9 kg/m² for age 18 years and older.

2.4. Statistical methods

Demographic and clinical variables were summarized by country, with categorical variables expressed as frequency and proportion, and continuous variables as means with standard deviation (SD) if parametric and medians with range if non-parametric. Differences between countries were compared using the Mann-Whitney test for continuous variables and the Chi-square test for categorical variables. Standardized mean difference (SMD) was calculated to identify statistically and clinically meaningful differences between the two countries, with a SMD of greater than 10 as the threshold [17]. The magnitude of difference in LF and poor nutrition between the two countries was estimated by linear regression and logistic regression, respectively. Confounders were chosen *a priori* based on clinical knowledge and prior literature and included: age, sex, genotype, diagnosis age, any *P. aeruginosa* infection, poor nutrition, and the pulmonary treatments recombinant human DNase (rhDNase) and macrolides. BMI was dichotomized as poor nutrition against the combined categories of adequate weight and overweight. Multiple imputation using chained equations was used to impute missing values for *P. aeruginosa* infection and pulmonary treatments. Ten imputed datasets were created and results were combined using Rubin’s rules [18]. Statistical analysis was conducted using R version 4.0.3. All p-values are two-sided and assessed at p < 0.05.

2.5. Ethical considerations

Approval has been obtained for the University of Cape Town Research Ethics Committee (HREC 032/2019) SA, and St. Michael’s Hospital (REB 20–310), Toronto. Informed consent and assents were collected in SA and Canada accordance with local and institutional and

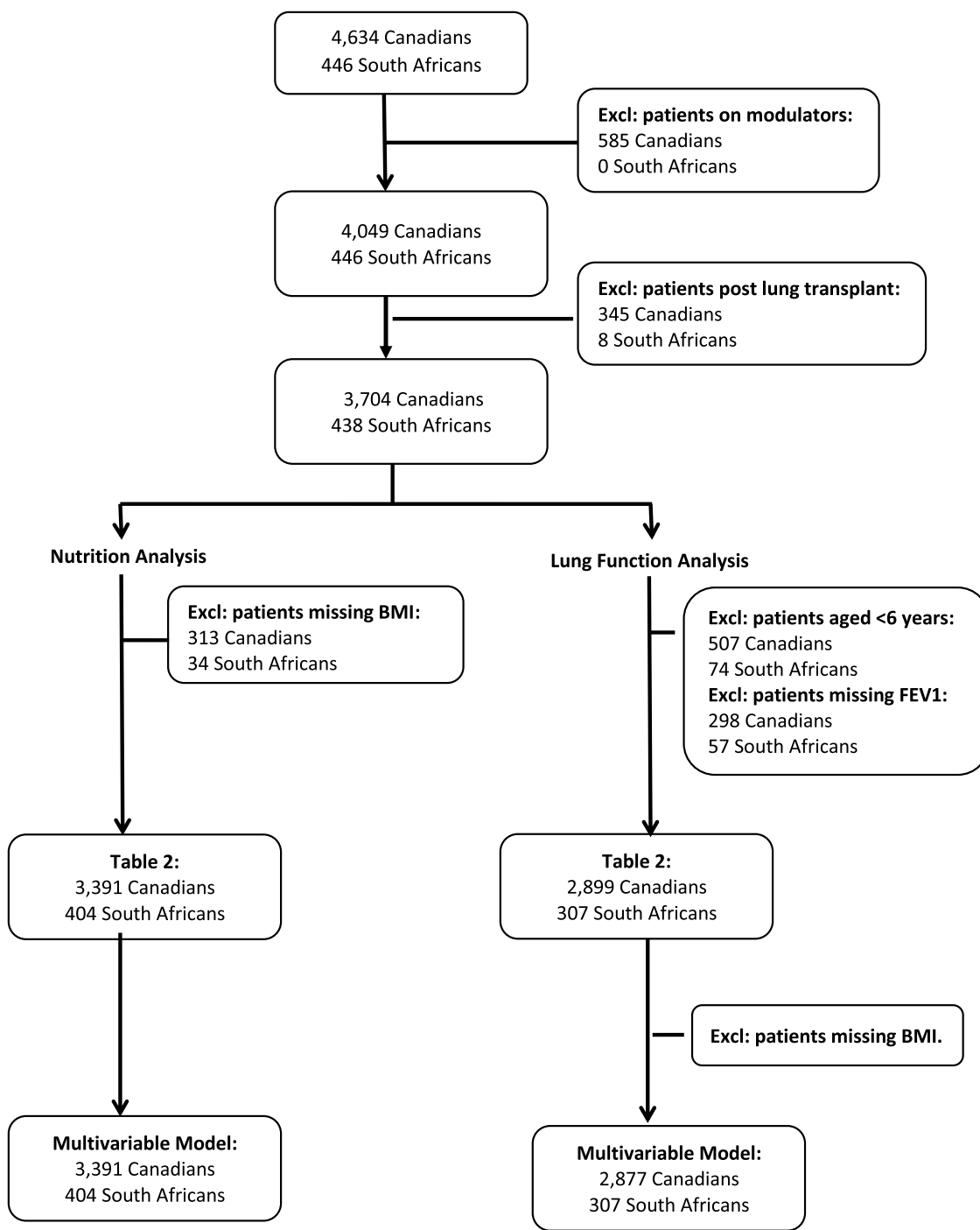


Fig. 1. Cohort flow diagram for Canadian and South African CF registry participant selection 01 January to 31 December 2018.

registry requirements.

3. Results

3.1. Demographic and diagnosis information

Data on 4049 Canadian and 446 SA pwCF in 2018 were analysed (Table 1 and Fig. 1). Compared to Canada, the CF population in SA was younger (median age 15.8 years vs. 24.1 years: SMD 52) and with a lower proportion of males (47.8% vs 54.2%; SMD 12.5) and White people (70.9% vs. 93.3%; SMD 61.3). Age of CF diagnosis was slightly younger in Canada and more children were diagnosed by NBS in Canada compared to SA ($n = 550, 13.6\%$ vs $n = 1, 0.2\%$, SMD 54.6); the one case

in SA was diagnosed by NBS in California, USA, and the family relocated to SA. Poor nutrition at time of diagnosis was present significantly more in SA compared to Canada ($n = 131, 53.5\%$ vs $n = 1117, 35.5\%$: SMD 36.8).

Sixty-one (1.3%) and 12 (2.7%) pwCF in Canada and SA, respectively, had incomplete genotyping recorded in the CF registries, Table 1. F508del was the most common CFTR mutation in both countries with similar proportions of people who were homozygous for F508del (SA 47.8% vs. Canada 42.4%). In SA, 3120+1G>A was the second most prevalent mutation (allele frequency 9.9%) and in Canada, 621+1G->T the second most prevalent mutation (allele frequency 2.0%). Overall, CFTR mutations class I-III were present with similar frequency in SA ($n = 384, 86.1\%$) and Canada ($n = 3426, 84.9\%$), Table 1.

Table 1

Demographic and diagnosis information among the CF population in Canada and South Africa, December 2018, excluding people on CFTR modulatory therapy.

	Canada (N = 4049)	South Africa (N = 446)	SMD
Sex: n (%)			
Female	1856 (45.8)	233 (52.2)	12.8
Male	2193 (54.2)	213 (47.8)	
Age in years, median (range)	24 (0.1–82.5)	15.8 (0–53.6)	52
Age in years: n (%)			
0–5.9 years	507 (12.5)	74 (16.6)	11.6
6–11.9 years	514 (12.7)	96 (21.5)	23.6
12–17.9 years	505 (12.5)	77 (17.3)	13.5
18–29.9 years	1061 (26.2)	121 (27.1)	2.1
≥ 30 years	1462 (36.1)	78 (17.5)	43
Race: n (%)			
White	3765 (93)	316 (70.9)	60
Mixed	28 (0.7)	85 (19.1)	64.7
Black African	34 (0.8)	41 (9.2)	39
Other/Unknown	222 (5.5)	4 (0.9)	26.3
Diagnosis age in years, median (range)	0.5 (0–75.8)	0.6 (0–47.8)	17.6
Diagnosis age in years: n (%)			
0–2	2696 (66.6)	307 (68.8)	4.8
2–18	1008 (24.9)	119 (26.7)	4.1
≥ 18	345 (8.5)	20 (4.5)	16.4
Newborn screening diagnosis: n (%)	550 (13.6)	1 (0.2)	54.6
Meconium ileus: n (%)	519 (12.8)	68 (15.2)	7.0
Nutritional status at diagnosis	n = 3146	n = 245	
Normal weight	1863 (59.2)	107 (43.7)	31.5
Poor Nutrition	1117 (35.5)	131 (53.5)	36.8
Overweight	166 (5.3)	7 (2.9)	12.3
Sweat test at diagnosis, mean (SD)			
sweat chloride test (mmol/L)	92.8 (23.8)	107.4 (21.9)	63.5
sweat conductivity (mmol/L)	N/A	105.8 (19.5)	N/A
no sweat test documented; n (%)	1879 (46.4)	152 (34.1)	25.3
Genotype: n (%)			
F508del:			
Homozygous	1715 (42.4)	213 (47.8)	10.9
Heterozygous	1774 (43.8)	145 (32.5)	23.4
Other	499 (12.3)	76 (17)	13.4
Missing /Incomplete genotyping (one or two unknown CFTR variants)	61 (1.5)	12 (2.7)	8.3
Mutation class: n (%)			
I – III	3426 (84.6)	384 (86.1)	7.8
IV- VI	502 (12.4)	44 (9.9)	
Missing/unclassified	121 (3)	18 (4)	5.7
Most common CFTR mutation allele frequencies: alleles, n (%)			
F508del	3436 (84.9)	358 (80.3)	N/A
3120+1G>A	0	44 (9.9)	
621+1G->T	80 (2.0)	0	
Unknown	65 (1.6)	12 (2.7)	
3272–26A>G	0	7 (1.6)	
Other	0	7 (1.6)	
G551D	0	3 (0.7)	
M1101K	41 (1.0)	0	
G542X	35 (0.9)	0	
711+1G->T	25 (0.6)	0	
A455E	21 (0.5)	5 (1.1)	
N1303K	21 (0.5)	5 (1.1)	
L206W	19 (0.5)	–	
3849+10kbC>T	0	1 (0.2)	
R1162X	0	1 (0.2)	
W1282X	15 (0.4)		

SMD: Standardized mean difference, values > 10 indicate clinically important differences.

3.2. Microbiology

P. aeruginosa was isolated in 2018 more frequently in SA than Canada (SA $n = 192$, 43.0% vs Canada $n = 1458$, 36.0%; SMD 26.3). Infection rates of MRSA and NTM were similar in both countries. Comparison of other infections or chronic infection status between countries could not be reliably determined due to different definitions and data capturing approaches.

3.3. Complications and CF therapies

The frequency of allergic bronchopulmonary aspergillosis (ABPA) and CF-related liver disease was similar in both countries, Table 2. CF-related diabetes (Canada 22.2% vs, SA 15.9%: SMD 12.7) and pancreatic sufficiency (Canada 18% vs, SA 10%; SMD 22) was more common in Canada than SA. Azithromycin and hypertonic saline inhalations were prescribed more frequently in SA and rhDNase more frequently used in Canada, Table 2. More pwCF in SA than Canada received one or more courses of intravenous antibiotics (SA $n = 202$, 45.3% vs Canada $n = 969$, 23.9%: SMD 46.4). Three hundred and seventy-six (9.3%) pwCF in Canada had received a lung transplant in 2018 or earlier compared to 11 (2.5%) in SA.

3.4. Outcomes

Forty-two (1.0%) and three (0.7%) adults in Canada and SA died in 2018, respectively (Table 2). There were no deaths in children less than 18 years in either country in 2018. Details of the number of LF and nutrition measurements in 2018 in pooled analysis is presented in Fig. 1.

3.4.1. Lung function

Lung function was significantly lower across all ages in SA compared to Canada, including children 6–11.9 years age (Fig. 2). After adjusting for age, sex, genotype, diagnosis age, any *P. aeruginosa* infection, poor nutrition, and pulmonary treatments, FEV_{1pp} was 8.9% lower in SA compared to Canada (95% CI 6.3% to 11.4%; p-value <0.0001), Table 3. The impact of nutrition on LF did not differ significantly by country.

3.4.2. Nutrition

Poor nutrition was significantly more prevalent across all age groups in SA compared to Canada, (SA 91, 22.5% vs Canada 373, 11.0%: SMD 31.2) including children 0–5.9 years, Table 2, and Fig. 3. After adjusting for age, sex, genotype, diagnosis age and any *P. aeruginosa* infection, poor nutrition was 1.7-fold more common in SA compared to Canada (OR 1.70; 95% CI 1.19–2.41). Other factors associated with poor nutrition in both countries were age of diagnosis 2–18 years, *P. aeruginosa* and pulmonary treatments, Table 3.

4. Discussion

This registry-based study comparing health outcomes between SA and Canada has identified significant health disparities, specifically less favorable LF and nutrition outcomes in SA compared to Canada that are attributable to factors over and above age, CFTR mutation class, sex, diagnosis age, *P. aeruginosa* infection, lung transplantation and CFTR modulator therapy. A further observation is that these differences were already present in early childhood suggesting early life health factors may be important determinants for poorer outcomes observed in SA. This study was not specifically designed to identify which additional factors were associated with poorer outcomes in SA. However, it is reasonable to assume that these may include lower SES, inferior standards of CF care, limited access to publicly funded health care and expensive therapies such as rhDNase and inhaled tobramycin. Socio-economic conditions and quality of healthcare delivery are important determinants of CF outcomes, even in high income settings [19,20]. We could not include robust measures of SES as these were not captured by either registry. However, the gross domestic product per capita of Canada is 7-fold higher than SA which indirectly reflects the difference in healthcare expenditure between the countries [21]. Although lung transplant recipients and pwCF on CFTR modulators were excluded from the formal analyses, inclusion of these factors would likely contribute to a wider difference in observed health outcomes between SA and Canada. Documenting and monitoring differences in CF-related outcomes is important in advocacy for improved quality of CF care and equitable global access to CFTR modulators [22].

Table 2

Clinical, lung function and nutritional characteristics of CF registry populations in South Africa and Canada in 2018, excluding people on CFTR modulatory therapy.

	Canada (N = 3391)	South Africa (N = 404)	SMD
Microbiology: n (%)			
Any <i>P. aeruginosa</i>	1458 (36)	192 (43)	26.3
Missing	157 (3.9)	65 (14.6)	37.6
Any MRSA	243 (6)	28 (6.3)	5.5
Missing	157 (3.9)	80 (17.9)	46.3
Any NTM isolate:	197 (4.9)	5 (1.1)	21.2
Missing	157 (3.9)	77 (17.3)	44.6
Pulmonary therapies: n (%)			
Hypertonic saline	1523 (37.6)	211 (50.6)	26.4
Recombinant human DNase	1904 (47)	122 (29.2)	37.4
Inhaled Antibiotics	2250 (55.6)	224 (53.8)	3.5
Azithromycin	1527 (37.7)	335 (80.3)	96.2
Intravenous antibiotic courses in 2018			
0	3080 (76.1)	244 (54.7)	46.1
1	579 (14.3)	101 (22.6)	21.6
2	224 (5.5)	52 (11.7)	22
≥ 3	166 (4.1)	49 (11)	26.3
lung transplant recipients (2018 or earlier)	376 (9.3)	11 (2.5)	28.9
liver transplant recipients (2018 or earlier)	28 (0.7)	0 (0)	N/A
Complications/comorbidity: n (%)			
ABPA	70 (1.8)	20 (5.2)	18.7
CF-related diabetes	900 (22.2)	71 (17.2)	12.7
CF-related liver disease with cirrhosis	135 (3.3)	22 (5.4)	10
Pancreatic insufficient	3310 (82.1)	377 (89.8)	22.1
Vital status in 2018			
Alive on 31 December 2018	4007 (99)	443 (99.3)	4
Died in 2018	42 (1)	3 (0.7)	4
Age at death (years): median (range)	33.7 (13.6–82.4)	36 (32–38)	7.4
Alive, but not seen in 2018	156 (3.9)	21 (4.7)	4.2
Nutritional status: n (%)			
Normal weight	2448 (72.2)	280 (69.3)	6.3
Overweight	570 (16.8)	33 (8.2)	26.4
Poor nutrition	373 (11.0)	91 (22.5)	31.2
Proportion with poor nutrition: n (%)			
F508del homozygous	186/1429 (13%)	38/194 (19.6%)	17.9
F508del heterozygous	137/1494 (9.2%)	32/133 (24.1%)	40.8
Class I-III mutation	328/2882 (11.4%)	79/353 (22.4%)	29.7
Class IV-V mutation	30/411 (7.3%)	7/35 (20%)	37.6
Lung function (age ≥ 6 years)			
n = 2899 (Canada) / n = 307 (South Africa)			
FEV₁pp: n (%)			
≥ 70	1962 (67.7)	171 (55.7)	24.8
40–69	705 (24.3)	101 (32.9)	19.1
< 40	232 (8)	35 (11.4)	11.5
FEV₁pp: Median (range)			
F508del homozygous	82.9 (61–98.7); N = 1296	78 (60.8–92.9); N = 180	15.6
F508del heterozygous	86.3 (61.3–100.2); N = 466	78 (53.1–92.7); N = 64	29.4
Other/missing genotype	84.9 (64.1–100.2); N = 1435	64.1 (51.7–85.8); N = 120	56.8
Class I-III mutation	83 (61.7–99.2); N = 2650	75.9 (56.2–90.1); N = 310	27.7
Class IV-V mutation	93 (71.1–103.2); N = 438	61.6 (51.4–85.1); N = 40	74.4

Legend Table 2:

NTM: non-tuberculous mycobacterium.

FEV₁z: Forced expiratory volume in one second z-score.

FEV₁pp: Forced expiratory volume in one second percent predicted.

ABPA: allergic bronchopulmonary aspergillosis.

MRSA: methicillin resistant *S. aureus*.

SMD: Standardised mean difference, values >10 indicate clinically important differences.

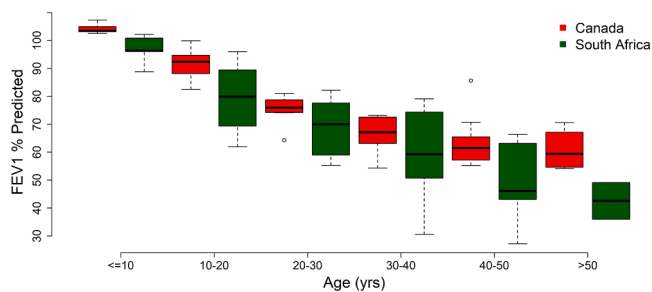


Fig. 2. Comparison of lung function (FEV₁pp) by age categories (6 years and older) between the South African and Canadian CF Registry cohorts in 2018. Horizontal bar represents median value and boxes 25–75% IQR; maximum and minimum range end indicated by thin horizontal bars.

Table 3

Adjusted multivariable analyses comparing lung function (age 6 years and older) and nutrition outcomes between South African (SA) and Canadian (CN) CF registry cohorts 2018.

Variable	Lung function (N = 2877 CN, 307 SA) [#]		Poor nutrition (N = 3391 CN, 404 SA) ^{#†}	
	Coefficient (95% CI)	P-value	Odds Ratio (95% CI)	P-value
Intercept	90.25 (88.48–92.03)	<0.0001	–	–
Country (SA vs CN)	–8.86 (–11.39–6.33)	<0.0001	1.70 (1.19–2.41)	0.0076
Sex (Male vs. female)	–1.15 (–2.59–0.30)	0.11	0.88 (0.70–1.11)	0.25
Mutation Class				
I-III	Ref		Ref	
IV- V	5.19 (2.88–7.49)	<0.0001	0.90 (0.58–1.39)	0.60
Missing	–2.20 (–6.13–1.73)	0.27	1.68 (0.92–3.05)	0.082
Age (centered at 18 years)				
*Age-18 – term 1	–685.09 (–733.20–636.98)	<0.0001	0.97 (0.96–0.98)	0.00012
*Age – 18 – term 2	221.31 (181.10–261.53)	<0.0001		
Age at diagnosis				
<2 years	Ref		Ref	
2–18 years	1.25 (–0.38–2.88)	0.13	1.25 (0.95–1.64)	0.10
≥ 18 years	11.02 (8.04–13.99)	<0.0001	1.09 (0.57–2.09)	0.78
Any <i>P. aeruginosa</i>				
	–2.88 (–4.47–1.30)	0.0010	1.21 (0.93–1.56)	0.14
Azithromycin				
	–9.27 (–10.96–7.58)	<0.0001	1.43 (1.06–1.91)	0.02
rhDNase				
	–7.14 (–8.68–5.61)	<0.0001	1.36 (1.05–1.75)	0.02
Poor nutrition				
	–16.16 (–18.30–14.02)	<0.0001	–	

Legend Table 3.

*Age was modeled as a quadratic relationship between age and lung function.

[#]Excluding lung transplant recipients, people started on CFTR modulator therapy before 2018 and children less than 6 years (no lung function recorded).

[†]Lung function excluded from the nutrition modeling to not exclude children less than 6 years without LF measurements.

A strength of this study is it illustrates the value of CF registries and how they can contribute to our understanding of CF epidemiology and outcomes in different global and socioeconomic settings. However, harmonization of registry variables as required in our study is key to meaningful comparisons between different registry populations [23].

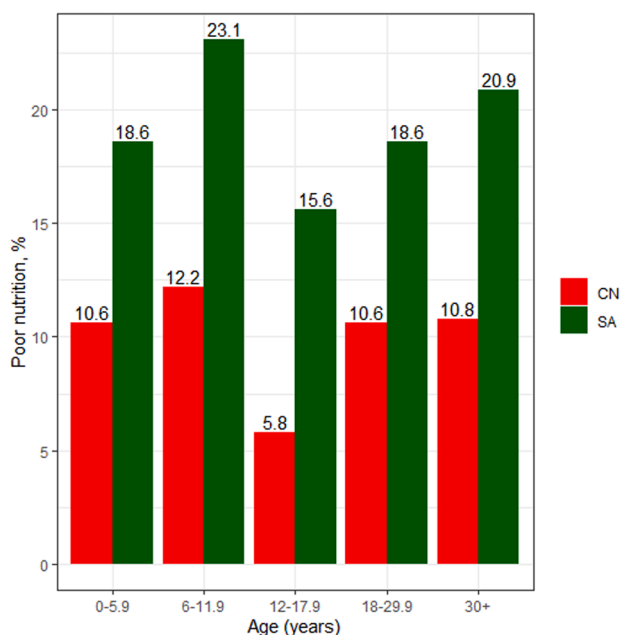


Fig. 3. Proportion (as percentage) of people with CF in the South African and Canadian CF Registry cohorts with poor nutrition in 2018, by age categories. Poor nutrition was defined as BMI z-score < -1 for age < 18 years or BMI < 18.5 kg/m² for age 18 years and older; differences between countries in all categories are statistically significant (SMD ≥ 10).

Our study is the first attempt to directly compare CF outcomes between a LMIC and HIC and highlights important disparities after adjusting for known factors and differences in genotype and ancestry characteristics between countries. South Africa shares similarities in CF epidemiology and unequal or limited health care resources with other LMIC. The disparity findings of our study are therefore relevant to other LMIC settings. Poor nutrition across all ages was more prevalent in SA compared to Canada and was the strongest factor negatively impacting LF in the adjusted multivariable analysis. Poor nutrition is an important modifiable aspect of CF care which needs greater attention in SA as similarly demonstrated in previous studies comparing nutrition outcomes between countries [23]. In addition, improved sputum surveillance and more aggressive *P. aeruginosa* treatment is another aspect of CF care which could be improved in SA.

People in LMIC represent 10% of an estimated 105,352 known diagnosed pwCF in 45 countries across the world of which only 12% are receiving triple combination CFTR modulator therapy [24]. However, the hidden global burden of undiagnosed CF disease is uncertain but estimated to be substantially greater than currently known and compounded by a paucity of high-quality data and limited capacity to diagnose CF in many LMIC [24]. Highlighting existing global disparities is especially important at a time when CFTR modulator therapy is rapidly becoming the standard of care in most HIC. Simulated projections of the Canadian CF Registry population have estimated that universal introduction of triple combination CFTR modulator therapy (tezacaftor/ivacaftor/elixacaftor) in 2021, would by 2030, reduce the number of CF-related deaths by 15% [4]. The delay and unlikely availability of CFTR modulator drugs in LMIC for a long time due to prohibitive pricing and patent protection laws, will undoubtedly lead to further widening of global disparities in CF outcomes between LMIC and HIC [25,26]. Ironically, earlier access to CFTR modulator therapy in LMIC will likely lessen the burden and complexity of CF care in LMIC due to less severe disease and fewer CF-related complications.

Another disparity highlighted by our study is the absence of NBS for CF in SA and the impact this likely has on explaining the lower nutritional status seen in the SA cohort. Although only 13% of Canadians

with CF in 2018 were diagnosed by NBS and median diagnosis age was slightly younger in Canada compared to SA, evidence that NBS improves long-term CF outcomes and survival is unequivocal [27]. Poorer nutrition and LF outcomes were present already in young SA children compared to children in Canada and supports the need to explore the feasibility of NBS for CF in SA. Although challenging and seldom prioritised by governments in LMIC, NBS for CF has been successfully implemented in several LMIC [27–30].

Our study has several limitations which were taken into consideration with interpretation of the findings. First, missing registry data significantly diminished the sample size of the formal analyses which reduced the precision of our estimates. Second, the SA registry data is unlikely to be an exhaustive representation of the CF population in SA as the true incidence, including undiagnosed cases of CF in SA, especially in the non-White population, is unknown. There is insufficient population data to accurately predict the expected incidence of CF in the majority, black African SA population but it is significantly lower than in White and mixed-race populations [31]. Data of the 446 SA patients in this study was extracted from the SA CF registry after its first year of inception in 2018 and is in our view the best representation of the available background diagnosed CF population in SA at the time. Follow-up and retention in the SA registry is high and it is unlikely that there are significant numbers of pwCF receiving care outside the designated CF clinics in SA, nor is it likely that CF-related deaths in SA registry patients were missed. Third, cross-sectional analysis of registry cohorts limit the interpretation of observed age-trends as they cannot be interpreted as average trends of individuals over time. Longitudinal analysis of individual trends over time between SA and Canada will enhance the interpretability of the observed differences observed in this study. Based on established knowledge, we selected variables *a priori* to include in the multivariable models for LF and nutrition outcomes. It is possible that inclusion of more variables such as a wider range of pulmonary infections might alter the overall differences in outcomes between countries due to unmeasured interactions. However, harmonization of all pulmonary infection definitions between the two registries was not possible, thus limiting the inclusion of other infections in the models. A slightly greater proportion of pancreatic sufficient pwCF in Canada compared to SA may be a factor contributing to better nutrition and LF outcomes in Canada. However, the proportional difference of people with pancreatic sufficiency is small and unlikely to be statistically significant. In addition, adjusting for class IV-VI CFTR mutations and older age of diagnosis would have accounted for the differences in proportions of pwCF with milder CF phenotypes with pancreatic sufficiency between the two countries.

In conclusion, this study has demonstrated the value of CF registry-based research and has highlighted important disparities in CF outcomes between a HIC and LMIC settings which has global relevance for countries with similar socioeconomic conditions and challenges impacting on CF care. In addition to improving the quality of established CF care, affordable and equitable global access to CFTR modulators in LMIC must be prioritised in order to reduce the inevitable widening CF outcome disparities between HIC and LMIC that will result from CFTR modulators becoming standard of care in HIC across the world.

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Declaration of Competing Interest

None.

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Supplementary materials

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